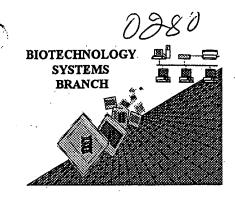
## RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following CRF diskette:

Application Serial Number:

09/458,813

Art Unit / Team No.:

OIPE

Date Processed by STIC:

12/30/99

THE ATTACHED PRINTOUT EXPLAINS THE ERRORS DETECTED.

PLEASE BE SURE TO FORWARD THIS INFORMATION TO THE APPLICANTS BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANTS ALONG WITH A NOTICE TO COMPLY or,
- 2) CALLING APPLICANTS AND FAXING THEM A COPY OF THE PRINTOUT WITH A NOTICE TO COMPLY

THIS WILL INSURE THAT THE NEXT SUBMISSION RECEIVED FROM THEM WILL BE ERROR FREE.

IF YOU HAVE ANY FURTHER QUESTIONS, PLEASE CALL:

**MARK SPENCER 703-308-4212** 

## Raw Sequence Listing Error Summary

## ERROR DETECTED SUGGESTED CORRECTION SERIAL NUMBER:

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE The number/text at the end of each line "wrapped" down to the next line. \_ Wrapped Nucleics This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3, as this will prevent "wrapping". The amino acid number/text at the end of each line "wrapped" down to the next line. Wrapped Aminos This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3, as this will prevent "wrapping". The rules require that a line not exceed 72 characters in length. This includes spaces. Incorrect Line Length The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs Misaligned Amino Acid between the numbering. It is recommended to delete any tabs and use spacing between the numbers. Numbering This file was not saved in ASCII (DOS) text, as required by the Sequence Rules. Non-ASCII Please ensure your subsequent submission is saved in ASCII text so that it can be processed. \_\_\_ contain n's or Xaa's which represented more than one residue. Variable Length As per the rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the (ix) feature section that some may be missing. A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid Patentin ver. 2.0 "bug" sequence(s) \_\_\_\_\_\_. Normally, Patentin would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. \_ missing. If intentional, please use the following format for each skipped sequence: Sequence(s) **Skipped Sequences** (2) INFORMATION FOR SEQ ID NO:X: (OLD RULES) (i) SEQUENCE CHARACTERISTICS:(Do not insert any headings under "SEQUENCE CHARACTERISTICS") (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: This sequence is intentionally skipped Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s). Sequence(s) \_\_\_\_ missing. If intentional, please use the following format for each skipped sequence. **Skipped Sequences** <210> sequence id number (NEW RULES) <400> sequence id number Use of n's and/or Xaa's have been detected in the Sequence Listing. Use of n's or Xaa's Use of <220> to <223> is MANDATORY if n's or Xaa's are present. (NEW RULES) In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents. \_\_\_ are missing this mandatory field or its response. Use of <213>Organism Sequence(s) \_\_\_ (NEW RULES) Sequence(s) \_\_\_\_ are missing the <220>Feature and associated headings. Use of <220>Feature Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial" or "Unknown" (NEW RULES) Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of new Rules) Please do not use "Copy to Disk" function of Patentin version 2.0. This causes a corrupted Patentin ver. 2.0 "bug" file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other means to copy file to floppy disk.

AKS-Biotechnology Systems Branch- 5/15/99

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<220> FEATURE:

## RAW SEQUENCE LISTING PATENT APPLICATION US/09/458,813

DATE: 12/30/1999

TIME: 14:42:54

Input Set: I458813.RAW

This Raw Listing contains the General Information Section and up to first 5 pages.

Corrected Diskette Needer Does Not Comply 1 <110> APPLICANT: Murphy, Brian R. 2 Collins, Peter L. 3 Durbin, Anna P. 4 Skiadopoulos, Mario H. 5 Tao, Tao <120> TITLE OF INVENTION: USE OF RECOMBINANT LIVE-ATTENUATED PARAINFLUENZA VIRUS 6 7 (PIV) AS A VECTOR TO PROTECT AGAINST DISEASE CAUSED BY 8 PIV AND RESPIRATORY SYNCYTIAL VIRUS (RSV) 9 <130> FILE REFERENCE: 17634-000330 10 <140> CURRENT APPLICATION NUMBER: US/09/458,813 11 <141> CURRENT FILING DATE: 1999-12-10 12 <150> EARLIER APPLICATION NUMBER: 09/083,793 13 <151> EARLIER FILING DATE: 1998-05-22 <150> EARLIER APPLICATION NUMBER: 60/047,575 <151> EARLIER FILING DATE: 1997-05-23 <150> EARLIER APPLICATION NUMBER: 60/059,385 <151> EARLIER FILING DATE: 1997-09-19 <160> NUMBER OF SEQ ID NOS: 30 18 19 <170> SOFTWARE: PatentIn Ver. 2.1 20 <210> SEQ ID NO 1 21 <211> LENGTH: 42 22 <212> TYPE: DNA 23 <213> ORGANISM: Artificial Sequence 24 <220> FEATURE: 25 <223> OTHER INFORMATION: Description of Artificial Sequence: Nucleotide 26 insert to conform inserted sequence to rule of 27 six. 28 <400> SEQUENCE: 1 29 cttaagaata tacaaataag aaaaacttag gattaaagag cg 42 30 <210> SEQ ID NO 2 <211> LENGTH: 36 31 <212> TYPE: DNA 33 <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Description of Artificial Sequence: Nucleotide 36 insert to conform inserted sequence to rule of 37 six. 38 <400> SEQUENCE: 2 39 gatccaacaa agaaacgaca ccgaacaaac cttaag 36 40 <210> SEQ ID NO 3 41 <211> LENGTH: 101 42 <212> TYPE: DNA 43 <213> ORGANISM: Artificial Sequence

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5

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VERIFICATION SUMMARY
PATENT APPLICATION US/09/458,813

DATE: 12/30/1999 TIME: 14:42:54

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